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A Novel Prodrug Strategy Based on Reversibly Degradable Guanidine Imides for High Oral Bioavailability and Prolonged Pharmacokinetics of Broad-Spectrum Anti-influenza Agents

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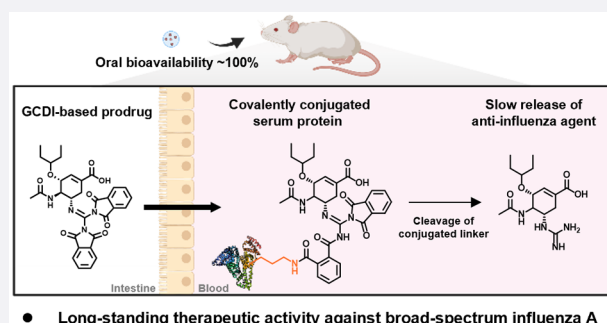
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ABSTRACT: We present orally administrable prodrugs (OSC-GCDIs) of guanidino oseltamivir carboxylate (GOC) based on guanidine cyclic diimide (GCDI) to treat influenza viruses. By concealing the guanidine group, which significantly limits the intestinal absorption, its prodrugs OSC-GCDIs demonstrate dramatic improvement of oral bioavailability. The most promising antiviral substance OSC-GCDI(P) readily forms covalent adducts with serum proteins via a degradable linker after the intestinal absorption. Subsequently, the active species, GOC, is released from the conjugate in a sustained manner, which greatly contributes to improving pharmacokinetic properties. Because of the remarkable improvements in both oral bioavailability and longevity of its active metabolite, OSC-GCDI(P) demonstrates outstanding therapeutic efficacy against both wild-type and oseltamivir-resistant (H275Y) influenza virus strains in a mouse infection model, even with a single oral administration.



INTRODUCTION

Oseltamivir phosphate (OS-P; Tamiflu), an ethyl ester prodrug of oseltamivir carboxylate (OSC), is used as a first-line therapy against both influenza A and B viruses, due to its potent antiviral efficacy and high oral bioavailability (>80%) (Figure 1a).^{1–5} However, the regimen for OS-P necessitates oral administration twice daily for 5 days, leading to inconvenience for patients. Furthermore, the emergence of OS-resistant strains, notably with the H275Y mutation in neuraminidase (NA), presents a great challenge and has become a global concern.^{6–13} These circumstances highlight the urgent need for novel therapeutics effective against wild-type and OS-resistant viruses, with reduced dosing frequency.

Among anti-influenza viral drug candidates, guanidino oseltamivir carboxylate (GOC) has been identified to be highly potent against wild-type and OS-resistant virus strains.^{1,5,14–16} The presence of the guanidine group provides additional interactions with the E226 residue in NA, resulting in a stronger binding affinity of GOC with the H275Y mutant NA when compared to OSC.^{11,16} However, GOC exhibits very low oral bioavailability (<4%), presenting a significant limitation in oral administration.^{5,16} Zanamivir, another guanidine-based antiviral agent, also faces challenges in oral administration and is consequently being developed as an inhalation drug.^{5,17}

Bioactive guanidine compounds often encounter oral bioavailability issues and are rarely found in orally admin-

istrable drugs. This difficulty mainly arises from the highly polar nature of the charged guanidine group, which impedes penetration through the intestinal barrier.^{18–21} To address these concerns, several approaches have been suggested, including conjugation with ligands,^{22–25} ion pairing,^{26–29} and acylation^{30–34} or *N*-hydroxylation^{16,35,36} of guanidine groups. However, each strategy has its own drawbacks, such as poorly enhanced oral bioavailability, compromised effectiveness, and complex synthetic routes.

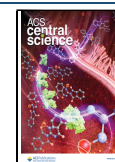
Previously, our group has reported a novel chemical structure termed guanidine cyclic diimides (GCDIs) (Figure 1b).³⁷ GCDI structures are easily synthesized through reactions between guanidines and cyclic anhydrides under mild conditions. Notably, the GCDI formation effectively conceals the positive charge of the free guanidine group, thereby enhancing lipophilicity. Moreover, GCDIs can undergo reversible degradation into the original guanidine species in the presence of various nucleophiles.

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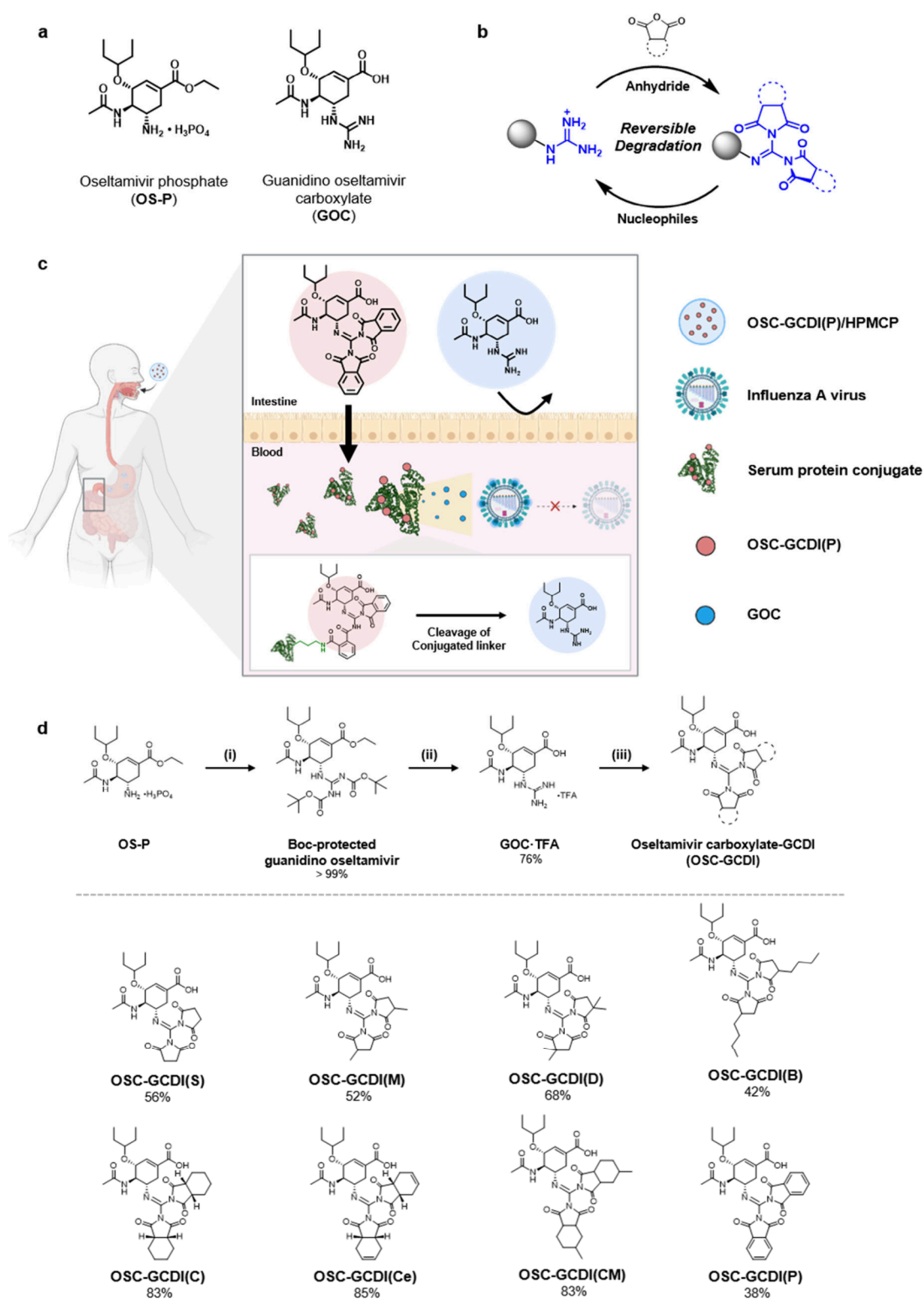


Figure 1. Novel prodrug strategy to enable oral administration of guanidine drugs utilizing the guanidine cyclic diimide (GCDI) structure. (a) Chemical structures of oseltamivir phosphate (OS-P) and guanidino oseltamivir carboxylate (GOC). (b) Reversible formation and degradation of GCDI. (c) Schematic illustration of the prodrug strategy to enhance oral bioavailability and pharmacokinetic half-lives of the guanidine drugs via installation of the GCDI moieties. (d) Synthesis and chemical structures of OSC-GCDIs.

On the basis of these findings, we anticipated that a novel prodrug strategy utilizing the GCDI structure could effectively address the issue of low oral bioavailability of guanidine drugs (Figure 1c). Because of a broad tolerance range of various functional groups under mild GCDI formation conditions,

GCDI moieties can be directly installed on guanidine groups without necessitating *de novo* synthetic pathways. Furthermore, by controlling the GCDI structures, the lipophilicity of prodrugs can be delicately tuned for facilitating efficient intestinal absorption. Once absorbed, the GCDI prodrugs can

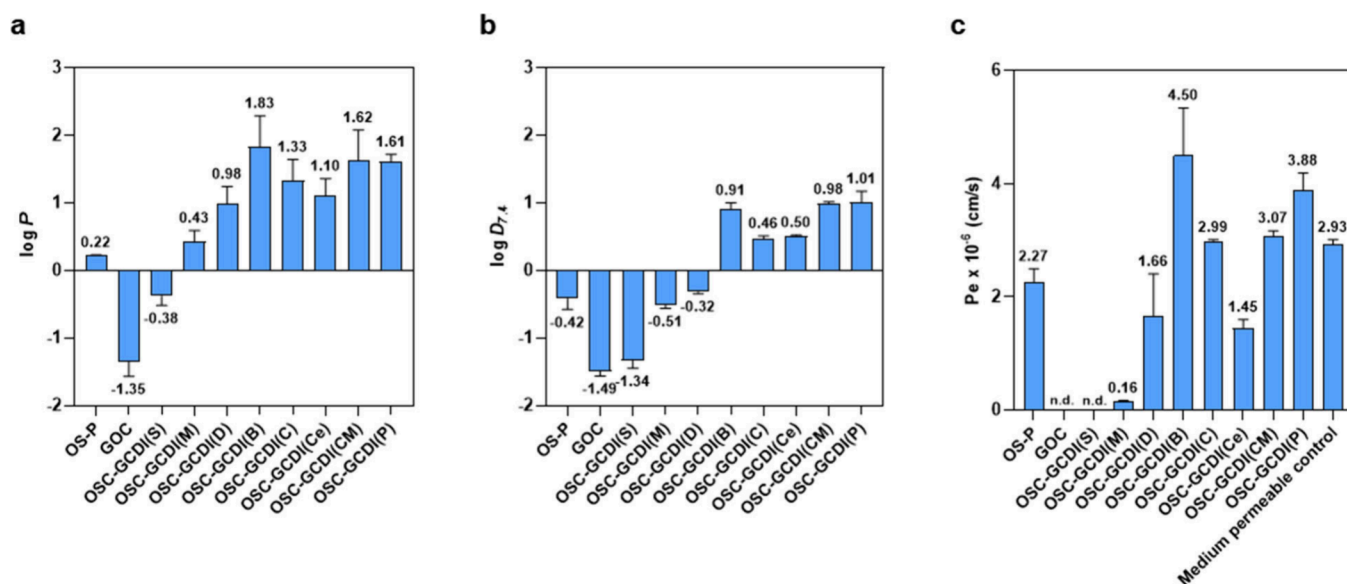


Figure 2. Lipophilicity and membrane permeability of OSC-GCDI prodrugs. (a) Octanol–water partition coefficients ($\log P$) of OS-P, GOC, and the OSC-GCDI prodrugs. (b) Octanol-phosphate buffer distribution coefficients at pH 7.4 ($\log D_{7.4}$) of OS-P, GOC, and the OSC-GCDI prodrugs. (c) Effective permeability through the artificial membrane of OS-P, GOC, and the OSC-GCDI prodrugs. The data are presented as means \pm standard deviations (SD) ($n = 3$). n.d., not detected.

be readily degraded in the bloodstream to release the active species.

In this work, we synthesized eight GCDI-based prodrugs of GOC, termed oseltamivir carboxylate guanidine cyclic diimides (OSC-GCDIs). Our results demonstrate that their enhanced lipophilicity significantly improved intestinal absorption *in vitro* and *in vivo*. More importantly, the absorbed OSC-GCDIs exhibited prolonged pharmacokinetic half-lives in the bloodstream, likely due to noncovalent or covalent interactions with serum proteins. The outstanding enhancement of both oral bioavailability and pharmacokinetics enabled OSC-GCDI(P) to exert excellent therapeutic effect against not only wild-type but also H275Y mutant influenza viruses in a mouse infection model, even with a single oral administration. We suggest OSC-GCDI(P) as a promising antiviral candidate against broad-spectrum influenza viruses. Moreover, the synthesis of GCDI-based prodrugs can serve as a robust platform to enhance the oral bioavailability and pharmacokinetics of various guanidine-based drugs.

RESULTS AND DISCUSSION

Synthesis of GCDI Prodrugs. The GCDI structure was introduced onto the guanidine group of GOC, an effective compound against the wild-type as well as H275Y mutant influenza viruses.^{1,5,14–16} Through reactions with various cyclic anhydrides (Figure 1d),³⁷ OSC-GCDIs were obtained in reasonable isolated yields (38–85%). To modulate the lipophilicity of the prodrugs, four different 5-membered cyclic anhydrides, succinic anhydride (S), methyl succinic anhydride (M), 2,2-dimethyl succinic anhydride (D), and *n*-butyl succinic anhydride (B), were reacted with GOC to form OSC-GCDI(S), OSC-GCDI(M), OSC-GCDI(D), and OSC-GCDI(B), respectively. Moreover, other four cyclic anhydrides with another 6-membered fused ring, *cis*-1,2-cyclohexanedicarboxylic anhydride (C), *cis*-1,2-cyclohexenedicarboxylic anhydride (Ce), 4-methyl-1,2-cyclohexenedicarboxylic anhydride (CM), and phthalic anhydride (P), were reacted with GOC to form OSC-GCDI(C), OSC-GCDI(Ce), OSC-

GCDI(CM), and OSC-GCDI(P), respectively. All the OSC-GCDIs could be directly synthesized from GOC, implying that the GCDI-based prodrug strategy can be generally applied to any guanidine-containing drugs at the final stages without requirement of *de novo* synthetic pathways for the prodrugs.

Lipophilicity of OSC-GCDIs. Lipophilicity is highly correlated to intestinal absorption rate and is regarded as a critical parameter in the development of orally administered drugs.³⁸ The octanol–water partition coefficient ($\log P$) and distribution coefficient ($\log D$) are common standards for assessing drug lipophilicity. Typically, $\log P$ and $\log D$ values at pH 7.4 ($\log D_{7.4}$) for orally administrable drugs range from -1 to 5 .^{39,40}

All the OSC-GCDIs showed significantly increased $\log P$ values compared to GOC, which exhibited low $\log P$ values of -1.35 (Figure 2a). In terms of $\log D_{7.4}$, which reflects the ionizable nature of the free carboxyl group, all the OSC-GCDIs, except for OSC-GCDI(S), showed notably increased $\log D_{7.4}$ values, ranging from -0.51 to 1.01 , compared to GOC with a $\log D_{7.4}$ value of -1.49 (Figure 2b). The lipophilicity of OSC-GCDIs well correlates with the GCDI structure, increasing with the imide hydrophobicity ($S < M < D < B$). OSC-GCDIs synthesized from fused-ring anhydrides showed $\log P$ and $\log D_{7.4}$ values of 1.10 – 1.62 and 0.46 – 1.01 , respectively. The lipophilicity of GOC increased from 8.8-fold in OSC-GCDI(S) to 2082-fold in OSC-GCDI(B) via the GCDI modification (based on P). Except for OSC-GCDI(S), the lipophilicity values of all OSC-GCDIs fell within the typical range for orally administrable drugs, illustrating the effectiveness of our GCDI prodrug strategy in enhancing lipophilicity. Notably, all the OSC-GCDIs except OSC-GCDI(S) and OSC-GCDI(M) exhibited even higher $\log P$ and $\log D_{7.4}$ values than OS-P, which has excellent oral bioavailability.

Parallel Artificial Membrane Permeability Assay (PAMPA) of OSC-GCDIs. While the precise absorption mechanism of the GCDI prodrugs remained unclear at that stage, it was plausible that these prodrugs would exhibit a

Table 1. Half-Lives of the OSC-GCDI Prodrugs in Phosphate Buffer and Serum

Compound	$T_{1/2}$ (min) ^a					
	Phosphate buffer				Serum	
	pH 2.0 ^b	pH 6.5 ^b	pH 7.4 ^b	pH 8.0 ^b	BALB/c ^c	Human ^c
OSC-GCDI(S)	>1440	770.2	101.9	42.0	17.5	16.7
OSC-GCDI(M)	>1440	866.4	108.3	39.5	13.5	18.4
OSC-GCDI(D)	>1440	>1440	495.1	161.2	158	117
OSC-GCDI(B)	>1440	>1440	462.1	157.5	17.1	52.5
OSC-GCDI(C)	>1440	630.1	81.6	26.8	23.8	26.2
OSC-GCDI(Ce)	>1440	1155	105.0	54.2	35.7	53.3
OSC-GCDI(CM)	>1440	693.1	99.0	3.92	13.3	31.2
OSC-GCDI(P)	93.7	87.7	14.1	<1	1.3	1.3

^aDetermined by HPLC analysis of the remaining prodrugs at various time points. The calculation of the half-lives was based on the assumption of pseudo-first-order kinetics of the degradation. ^bExamined in phosphate-buffered saline (50 mM phosphate, 154 mM ionic strength, 37 °C). ^cExamined in serum at a prodrug concentration of 200 μ M at 37 °C.

higher passive diffusion rate through the duodenal mucosa compared to GOC. To assess this hypothesis, we conducted the parallel artificial membrane permeability assay (PAMPA) for simulating passive transcellular permeation.⁴¹

The permeation rate of each OSC-GCDI through the artificial membrane was calculated by measuring the effective permeability (P_e). The permeability of GOC was almost negligible under these conditions; however, all OSC-GCDIs except OSC-GCDI(S) exhibited a remarkable increase in P_e (Figure 2c). In line with the lipophilicity tendency, OSC-GCDIs with bulkier imide groups tended to have higher P_e values. It is noteworthy that four OSC-GCDIs, specifically OSC-GCDI(B), OSC-GCDI(C), OSC-GCDI(CM), and OSC-GCDI(P), demonstrated P_e values that were comparable to or even exceeding those of OS-P ($P_e = 2.27 \times 10^{-6}$ cm/s) and a control compound with medium permeability ($P_e = 2.93 \times 10^{-6}$ cm/s). These findings led us to expect that GCDI derivatization could significantly enhance the intestinal absorption of guanidine drugs, potentially enabling their successful oral administration.

Activation of OSC-GCDIs and Conjugation to Albumin. Our group has previously demonstrated that the GCDI structures are readily degraded, releasing their parent guanidine molecules in the presence of various nucleophiles.³⁷ Consequently, we anticipated that GCDI prodrugs would regenerate the active drug, GOC, after intestinal absorption. Initially, we evaluated the stability or degradability of OSC-GCDIs (Table 1 and Figure S1) in phosphate buffers with different pHs. In general, OSC-GCDIs featuring bulkier imide groups exhibited greater stability against hydrolysis, likely due to reduced water accessibility to the imide carbonyl groups. Moreover, the hydrolysis rate of the GCDI prodrugs escalated with increasing pH. At pH 2.0 (mimicking stomach conditions), all OSC-GCDIs, except OSC-GCDI(P) with a half-life of 94 min, maintained excellent stability with half-lives longer than 24 h. At pH 6.5 (representing duodenal conditions), most prodrugs presented half-lives beyond 5 h, except for OSC-GCDI(P) with a half-life of 88 min. The least stable OSC-GCDI(P) exhibited half-life of 14.1 min at pH 7.4, whereas the most stable OSC-GCDI(D) demonstrated a half-life exceeding 8 h at pH 7.4. Other prodrugs showed half-lives ranging from 1 to 8 h at pH 7.4, and from 3 min to 3 h at pH 8.0. As suggested in previous studies,⁴² we suspect that the exceptional hydrolysis rate of OSC-GCDI(P) can be attributed to additional ring strain of the phthalimide structure to maximize the electron delocalization, which may lead to

intrinsic instability of OSC-GCDI(P). Additionally, given that aromatic carboxylic acids generally have higher acidity than aliphatic acids, the negative charge of the amic acid intermediate, which is generated by the degradation of OSC-GCDI(P), is expected to be stabilized more efficiently in the aromatic ring compared to those from other OSC-GCDIs with the aliphatic imide structures. These characteristics may contribute to the faster hydrolysis of OSC-GCDI(P).

To ascertain if OSC-GCDIs convert into their active form, GOC, after absorption into the bloodstream, we incubated the OSC-GCDIs in BALB/c mouse and human serum and then quantified both unhydrolyzed OSC-GCDIs and hydrolyzed GOC (Table 1 and Figure S2). OSC-GCDIs underwent significantly faster degradation in serum than in phosphate buffers. Most OSC-GCDIs showed half-lives of approximately 10–30 min, with bulkier imide groups generally associated with longer half-lives. OSC-GCDI(P) displayed the shortest half-life (1.3 min), while OSC-GCDI(D) had the longest (>100 min) in serum. The degradation rate of OSC-GCDIs was not accelerated by Sprague–Dawley rat liver microsomes (Table S1). The results supported that the degradation of OSC-GCDIs or activation to GOC might be based primarily on reactions in serum, not by hepatic metabolism.

The liberation of GOC was notably slower than the degradation of GCDI species in serum (Figure S2). Particularly, OSC-GCDI(P), which exhibited only a half-life of 1.3 min in serum, released 37% of GOC after 30 min and 68% after 24 h. Similarly, OSC-GCDI(M) and OSC-GCDI(C) released GOC in a sustained and gradual manner after the original compounds have completely disappeared, suggesting the existence of intermediates prior to the full hydrolysis into GOC. Given their structural features, the conversion of OSC-GCDIs to GOC theoretically requires four nucleophilic attacks on the imide rings, possibly yielding various guanidine amic acids as likely intermediates (Figure S3).

The more rapid degradation of OSC-GCDIs in serum compared to buffers implies the possibility of nucleophilic attacks on the GCDI structure by nonwater nucleophiles. We suspected that the nucleophilic residues of serum proteins could attack the GCDI imide rings. The increased lipophilicity of OSC-GCDIs might facilitate their noncovalent association with albumin, thereby increasing the probability of nucleophilic attacks on the GCDI structures. Corroborating this hypothesis, MALDI-TOF MS analysis of the mixture of OSC-GCDI(P) and human serum albumin (HSA) provided the

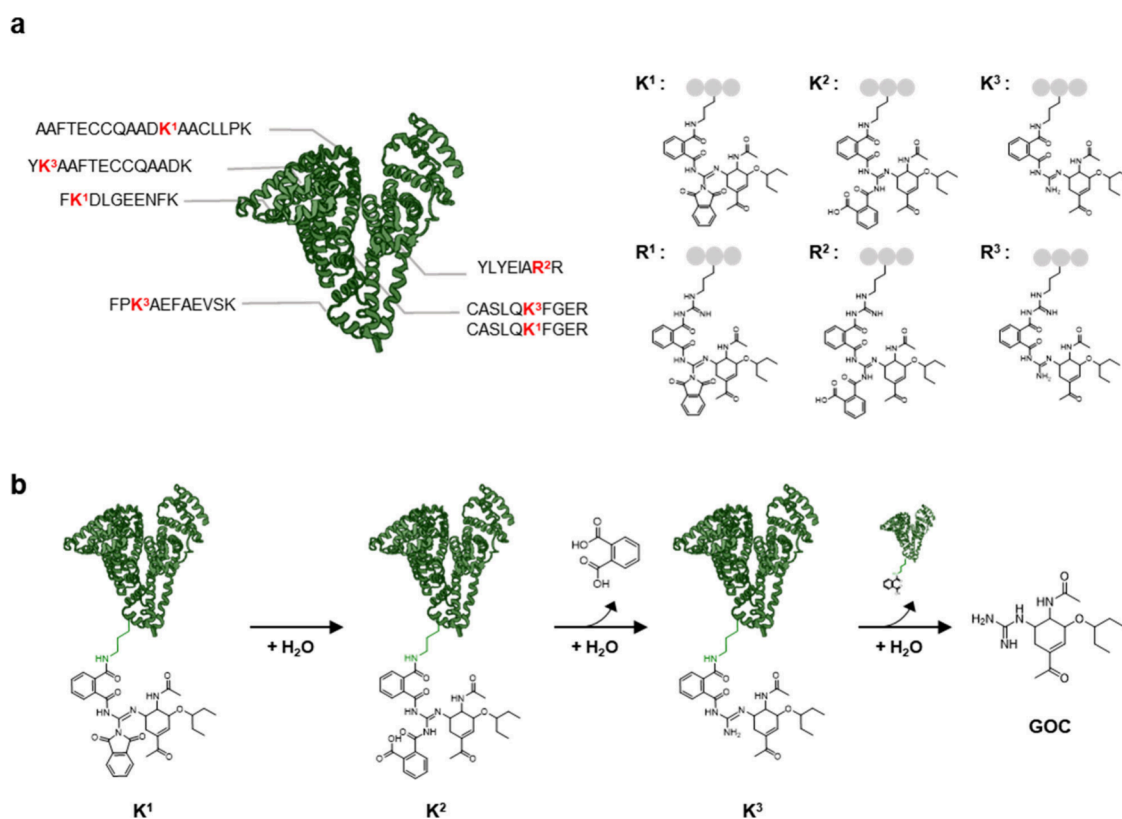


Figure 3. Covalent conjugation of OSC-GCDIs to human serum albumin (HSA). (a) Amino acid residues conjugated with OSC-GCDI in HSA, which were identified by MALDI-TOF MS. Suggested structures of the covalent linkage between the sequence and OSC-GCDI(P) are shown. (b) A proposed mechanism of the GOC release from HSA. The lysine adduct of OSC-GCDI(P), K¹, can be successively hydrolyzed to K², K³, and finally, GOC.

evidence of covalent adducts between them (Figures 3a and S4). Several lysine and arginine residues in HSA were covalently conjugated to the prodrug via amide or acyl guanidine linkages.

The covalent conjugation of the GCDI prodrugs onto serum proteins can provide a new strategy to increase pharmacokinetic half-lives. Similar to other albumin-conjugated drugs,^{43–46} the GCDI prodrugs are likely to remain in the bloodstream in their conjugated form with albumin, thereby reducing their hepatic intracellular metabolism and excretion rates. Considering the vulnerability of the guanidine amide linkage between GOC and albumin to nucleophilic attacks, it was postulated that GOC could be gradually released from albumin via hydrolytic intermediates (Figure 3b). Indeed, incubation of OSC-GCDI(P)-HSA and OSC-GCDI(C)-HSA conjugates resulted in the release of GOC in a sustained manner for at least 72 h (Figure S5).

In Vitro Antiviral Activity of OSC-GCDIs. To evaluate the antiviral effectiveness of OSC-GCDIs and GOC, we conducted a cell-based assay to assess the cytopathic effects resulting from the infection of different influenza virus (sub)types, such as A/Puerto Rico/8/1934 (PR8; H1N1), A/Hong Kong/8/1968 (HK; H3N2), and B/Lee/1940 (Lee) (Table 2). OSC-GCDIs exhibited comparable activity against PR8, with EC₅₀ values ranging from 0.01 to 0.09 μM. For HK-infected cells, the EC₅₀ values was below 0.04 μM, with OSC-GCDI(C) showing the most potent inhibition (EC₅₀, < 0.005 μM). These compounds demonstrated variable antiviral efficacy against the influenza B virus strain, Lee, with EC₅₀ values ranging from 2.21 to 72.38 μM, and the lowest values

observed for OSC-GCDI(C) and OSC-GCDI(P). Collectively, among the OSC-GCDIs tested, OSC-GCDI(C) and OSC-GCDI(P) demonstrated relatively more potent antiviral activity against influenza A and B viruses, *in vitro*.

To evaluate broad-spectrum antiviral efficacy, particularly against OS-resistant viruses with the H275Y mutation in NA, we further evaluated the change in antiviral effectiveness against the reverse genetically rescued PR8 virus (rgPR8) and its corresponding H275Y mutant strain (rgPR8(H275Y)) (Table 2). The calculated resistance factors (RFs), defined as the ratio of EC₅₀ values against rgPR8(H275Y) to wild-type rgPR8, indicated a substantial reduction in antiviral resistance for all OSC-GCDIs (RFs, 14.1–190) and GOC (RF, 18.0), both of which were discriminated from that of OSC (RF, 274). Additionally, we investigated the antiviral activity of these OSC-GCDIs against other H275Y mutants, A/Korea/2785/2009 (KR2785) and rgA/Korea/09/2009Δ53–60 (rgKR09). The OSC-GCDIs surpassed OSC in antiviral activity against the H275Y mutant viruses, although they were not as potent as GOC. Given that full activation from OSC-GCDIs to GOC take a substantial amount of time in buffers or serum (Table 1 and Figures S1–S3), the observed reduced *in vitro* efficacy of the GCDI prodrugs is understandable. Among the OSC-GCDIs, OSC-GCDI(S), OSC-GCDI(M), OSC-GCDI(C), and OSC-GCDI(P) exhibited superior activity relative to the remaining four compounds. Taken together, GCDI formation from OSC appeared to contribute to recovering its reduced antiviral activity against the H275Y mutant viruses.

In Vivo Pharmacokinetics and Antiviral Activity of OSC-GCDIs. GOC is more potent than OSC both against

Table 2. Antiviral Activity of OSC-GCDI Prodrugs against Influenza A and B Viruses in MDCK Cells

Compound	CC ₅₀ (μ M) ^a	EC ₅₀ (μ M) ^b (S.I.) ^c						
		PR8 ^d	HK ^e	Lee ^f	rgPR8 ^g	rgPR8 (H275Y) ^h	KR2785 (H275Y) ⁱ	rgKR09 (H275Y) ^j
OSC-GCDI(S)	>100	0.06 \pm 0.01 (>1667)	0.03 \pm 0.01 (>3333)	11.10 \pm 1.20 (>9.01)	0.07 \pm 0.01 (>1429)	0.99 \pm 0.01 (>101)	8.55 \pm 2.15 (>11.7)	2.20 \pm 0.10 (>45.5)
OSC-GCDI(M)	>100	0.04 \pm 0.01 (>2500)	0.02 \pm 0.01 (>5000)	5.55 \pm 1.65 (>18.0)	0.02 \pm 0.01 (>5000)	0.74 \pm 0.05 (>135.1)	4.61 \pm 0.35 (>21.7)	1.96 \pm 0.15 (>51.0)
OSC-GCDI(D)	>100	0.07 \pm 0.01 (>1429)	0.03 \pm 0.01 (>3333)	13.70 \pm 2.85 (>7.3)	0.05 \pm 0.01 (>2000)	1.52 \pm 0.20 (>65.8)	9.25 \pm 2.40 (>10.8)	4.37 \pm 1.10 (>22.9)
OSC-GCDI(B)	>100	0.09 \pm 0.01 (>1111)	0.04 \pm 0.01 (>2500)	72.38 \pm 1.35 (>1.4)	0.08 \pm 0.01 (>1250)	3.41 \pm 0.01 (>29.3)	17.59 \pm 2.90 (>5.7)	12.86 \pm 0.65 (>7.8)
OSC-GCDI(C)	>100	0.01 \pm 0.01 (>10,000)	<0.005 (>20,000)	2.21 \pm 0.20 (>45.2)	0.01 \pm 0.01 (>10,000)	0.49 \pm 0.01 (>204.1)	5.91 \pm 0.85 (>16.9)	3.70 \pm 0.50 (>27.0)
OSC-GCDI(Ce)	>100	0.03 \pm 0.01 (>3333)	0.02 \pm 0.01 (>5000)	11.96 \pm 4.85 (>8.4)	0.01 \pm 0.01 (>10,000)	1.90 \pm 0.10 (>52.6)	13.22 \pm 0.10 (>7.6)	5.21 \pm 0.02 (>19.2)
OSC-GCDI(CM)	>100	0.03 \pm 0.01 (>3333)	0.01 \pm 0.01 (>10,000)	8.36 \pm 1.00 (>12.0)	0.01 \pm 0.01 (>10,000)	1.20 \pm 0.05 (>83.3)	13.98 \pm 5.15 (>7.2)	4.43 \pm 0.25 (>22.7)
OSC-GCDI(P)	>100	0.03 \pm 0.01 (>3333)	0.01 \pm 0.01 (>10,000)	2.26 \pm 0.40 (>44.2)	0.01 \pm 0.01 (>16,667)	0.48 \pm 0.01 (>208.3)	4.61 \pm 0.35 (>21.7)	4.61 \pm 0.05 (>21.7)
GOC	>100	0.005 \pm 0.001 (>20,000)	<0.005 (>20,000)	0.70 \pm 0.15 (>142.9)	0.005 \pm 0.001 (20,000)	0.09 \pm 0.01 (>1111)	0.78 \pm 0.01 (>128.2)	0.11 \pm 0.01 (>909.1)
OSC	>100	0.10 \pm 0.01 (>1000)	<0.005 (>20,000)	1.29 \pm 0.25 (>77.5)	0.05 \pm 0.01 (>2000)	13.71 \pm 2.70 (>7.3)	56.61 \pm 0.30 (>1.8)	25.87 \pm 0.65 (>3.9)
RBV ^k	>100	11.1 \pm 0.01 (>9.0)	21.78 \pm 1.10 (>4.6)	13.41 \pm 0.95 (>7.5)	15.95 \pm 1.55 (>6.3)	16.43 \pm 0.45 (>6.1)	18.26 \pm 1.30 (>6.6)	15.79 \pm 0.15 (>6.3)
AMT ^l	>100	>100 (n.d.) ^m	1.56 \pm 0.40 (>64.1)	>100 (n.d.)	>100 (n.d.)	>100 (n.d.)	>100 (n.d.)	>100 (n.d.)

^aFifty percentage cytotoxic concentration to MDCK cells. ^bFifty percent effective concentration against influenza virus. ^cSelectivity index, the ratio of CC₅₀ to EC₅₀. ^dA/Puerto Rico/8/1934 (H1N1). ^eA/Hong Kong/8/1968 (H3N2). ^fB/Lee/1940. ^gReverse genetically generated A/Puerto Rico/8/1934. ^hReverse genetically generated A/Puerto Rico/8/1934 with the H275Y mutation in NA. ⁱA/Korea/2785/2009 (H1N1) with the H275Y mutation in NA. ^jReverse genetically generated A/Korea/09/2009 Δ 53–60 with the H275Y mutation and amino acid deletion between 53 and 60 in NA. ^kRibavirin. ^lAmantadine. ^mNot determined. The data are presented as means \pm standard deviations (SD) ($n = 3$).

wild-type and OS-resistant influenza viruses (Table 2).^{16,47} However, the limited oral bioavailability of GOC has hampered its effectiveness for clinical development. Encouraged by the improved *in vitro* permeability of OSC-GCDIs and their successful conversion to active GOC in serum, we explored the *in vivo* pharmacokinetic (PK) properties and antiviral activity of OSC-GCDI(C) and OSC-GCDI(P), which exhibited the most potent and broad-spectrum antiviral activity *in vitro* (Table 2).

In preliminary experiments, GOC as a trifluoroacetate salt (GOC·TFA), OSC-GCDI(C), and OSC-GCDI(P) were orally administered to mice. Pharmacokinetic analysis over 24 h revealed that GOC·TFA had an oral bioavailability (F_t) of 9.69% and a plasma half-life ($T_{1/2}$) of 4 h (Figure S6a). However, the F_t values for OSC-GCDI(C) and OSC-GCDI(P) were 0.48% and 5.95%, respectively (Figure S7), less than that of GOC·TFA. Notably, we found that OSC-GCDI(P) exhibits a remarkably longer $T_{1/2}$ (6.93 h) compared to GOC·TFA, which has a $T_{1/2}$ of 4.00 h, and OSC-GCDI(C), for which $T_{1/2}$ could not be estimated. These results suggest that although orally administered OSC-GCDI(P) has limited bioavailability, it facilitates sustained release of the active metabolite, GOC, in mouse blood.

On the basis of the extended half-life of OSC-GCDI(P), we hypothesized that consistent concentrations of GOC metabolized from orally administered OSC-GCDI(P) could effectively protect mice from influenza virus infection. Nevertheless, the therapeutic efficacy of OSC-GCDI(P) in a

mouse infection model was inferior to those of OS-P and GOC, as measured by dose response in survival rates and mortality (Figure S8), likely attributed to suboptimal C_{max} and resulting insufficient oral bioavailability. We assumed that these relatively poor antiviral activity of OSC-GCDI(P) in oral administration especially at lower doses could result from its vulnerability against nucleophiles prior to intestinal adsorption. Although OSC-GCDI(P) is stable in acidic buffers simulating stomach pH, it is prone to hydrolysis into GOC under neutral or basic conditions (Table 1), and to covalent binding with proteins by their nucleophilic attack to OSC-GCDI(P) along the gastrointestinal track (Figure 3). Such reactions may result in the accumulation of metabolites or metabolic intermediates from OSC-GCDI(P) with reduced intestinal absorption, underscoring the need of formulation for protecting it from premature degradation.

***In Vivo* Pharmacokinetics and Antiviral Activity of OSC-GCDIs Formulated with HPMCP.** To minimize the gastrointestinal degradation before intestinal absorption, OSC-GCDIs were formulated with a gastro-retentive drug delivery vehicle, hydroxypropyl methylcellulose phthalate (HPMCP), designated as OSC-GCDI/HPMCP. OSC-GCDI(C) and OSC-GCDI(P) were successfully encapsulated within HPMCP through coprecipitation in an acidic solution (Figure S9). The HPMCP encapsulation itself has no significant effect on the passive diffusion into membranes as shown in the PAMPA results of GOC/HPMCP and OSC-GCDI(P)/HPMCP (Table S2).

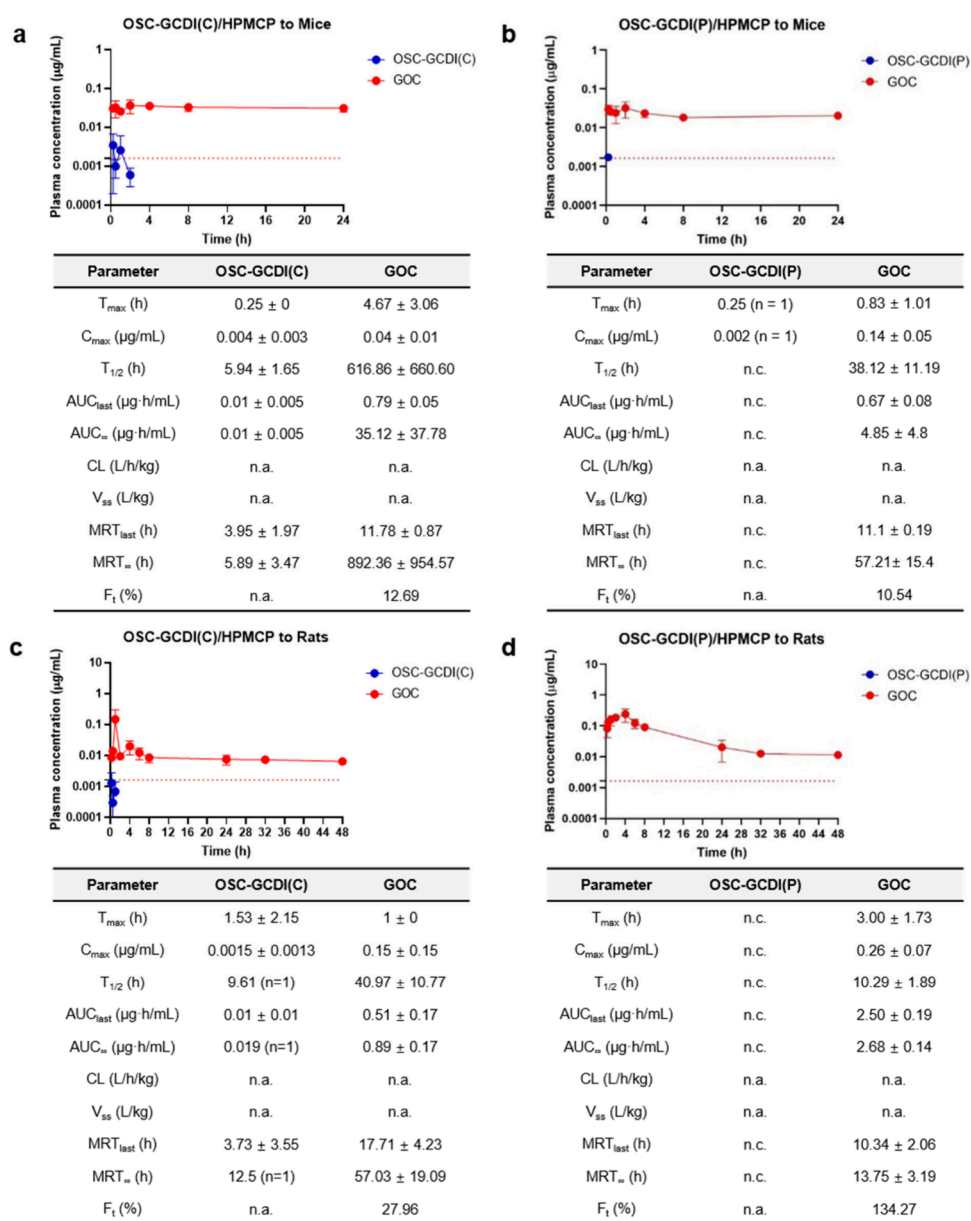


Figure 4. Pharmacokinetic properties of OSC-GCDIs/HPMCP in mice and rats. (a, b) Plasma concentrations and pharmacokinetic parameters of OSC-GCDIs (blue ●) and GOC (red ●) at various time points after oral administration of (a) OSC-GCDI(C)/HPMCP and (b) OSC-GCDI(P)/HPMCP to mice at doses of 5 mg/kg based on the amount of OSC-GCDIs within the complex. (c, d) Plasma concentrations and pharmacokinetic parameters of OSC-GCDIs (blue ●) and GOC (red ●) at various time points after oral administration of (c) OSC-GCDI(C)/HPMCP and (d) OSC-GCDI(P)/HPMCP to rats at doses of 10 mg/kg based on the amount of OSC-GCDIs within the complex. The data are presented as means ± standard deviations (SD) from three mice or rats, except for those from OSC-GCDI(C) in panel (a) at 0.25 h and OSC-GCDI(P) in panel (b) at 0.25 h, which were detected in one mouse among the three. The GOC concentration in plasma equivalent to an EC_{50} value (0.005 µM) against PR8 (A/H1N1) in MDCK cells is indicated by red dashed lines. n.a., not applicable; n.c., not calculated.

In an initial mouse pharmacokinetics analysis for 24 h, the F_t values of OSC-GCDI(C)/HPMCP and OSC-GCDI(P)/HPMCP were determined to be 12.69% and 10.54%, respectively, which were notably higher than those of unformulated OSC-GCDI(C) and OSC-GCDI(P) (Figure 4a and b, and Figure S7a and b). More importantly, we found that plasma GOC concentrations were almost constant even 24 h postadministration, implying the possibility of prolonged GOC release in the bloodstream. Consequently, we extended our pharmacokinetic analysis to rats for a longer period of 48 h, observing that OSC-GCDI(C)/HPMCP and OSC-GCDI(P)/HPMCP exhibited remarkably higher F_t values of 27.96% and

134.27%, respectively (Figure 4c and d), compared to GOC-TFA with an F_t value of 2.36% (Figure S6b). The F_t exceeding 100% for OSC-GCDI(P)/HPMCP may be attributed to the slower clearance rate of OSC-GCDI(P) as a result of covalent interactions with plasma proteins.^{48,49} These findings proved enhanced intestinal adsorption of OSC-GCDIs when encapsulated with HPMCP. In addition, given that the P_e value of OSC-GCDI(C) is comparable to that of OSC-GCDI(P) (Figure 2c), we suspect that the lower F_t value of OSC-GCDI(C) compared to that of OSC-GCDI(P) probably did not result from poor intestinal absorption of OSC-GCDI(C). Considering that F_t is defined as the molar ratio of detected

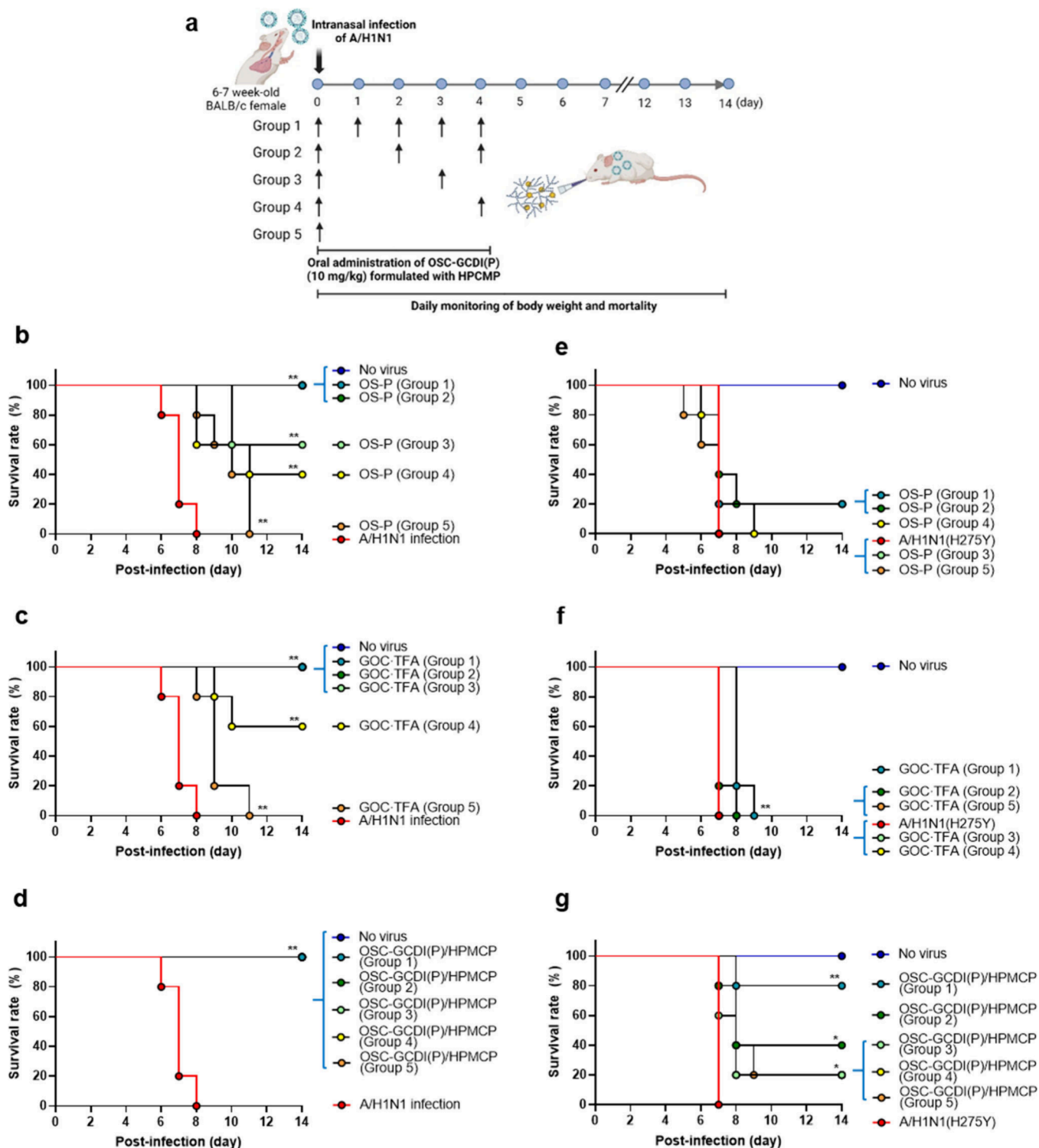


Figure 5. Effect of oral administration frequency on therapeutic efficacy of OSC-GCDI(P)/HPMCP in mice infected with a wild-type influenza virus and an OS-resistant influenza virus harboring the H275Y mutation in NA. (a) Schematic representation illustrating the administration frequency of OSC-GCDI(P) formulated with HPCMP to mice infected by wild-type mouse-adapted virus (maPR8) or the OS-resistant A/H1N1 virus (rgA/Korea/09/2009 Δ 53–60(H275Y)), referred to as A/H1N1(H275Y). Each compound was additionally administered at different intervals: days 1 to 4 postinfection in group 1, days 2 and 4 in group 2, day 3 in group 3, and day 4 in group 4. In Group 5, there was no additional treatment after the single administration. (b–d) Survival rates of maPR8-infected mice after (b) OS-P, (c) GOC-TFA, or (d) OSC-GCDI(P) administration. (e–g) Survival rates of A/H1N1(H275Y)-infected mice after (e) OS-P, (f) GOC-TFA, or (g) OSC-GCDI(P) administration. Groups with overlapping survival rate curve are indicated with blue brackets. Statistical significance was determined by comparing time-course survival rates to the maPR8- or A/H1N1(H275Y)-infected group. *, $P < 0.05$; **, $P < 0.01$.

active species (GOC) to orally administered prodrug (OSC-GCDI), not only intestinal absorption but also activation rate

in vivo is a crucial factor. Therefore, the lower F_t of OSC-GCDI(C) may be due to slower activation of OSC-GCDI(C)

into GOC in the bloodstream (Figures S2 and S5), which may lead to incomplete activation within 48 h in the rat model. Moreover, OSC-GCDI(C)/HPMCP and OSC-GCDI(P)/HPMCP exhibited $T_{1/2}$ of 40.97 and 10.29 h, respectively, which were significantly longer than that of GOC·TFA (3.57 h). The four times longer $T_{1/2}$ of OSC-GCDI(C) than that of OSC-GCDI(P) also supports our argument for the slower activation of OSC-GCDI(C) than OSC-GCDI(P). Furthermore, OSC-GCDI/HPMCP formulations maintained the GOC concentrations above the EC_{50} value (5.0 nM equal to 1.63 ng/mL) for at least 2 days. The higher lipophilicity of OSC-GCDIs could facilitate noncovalent binding to albumin, thereby prolonging circulation.^{50–52} Taken together, the result suggested that OSC-GCDIs circulates as a covalently conjugated form with serum proteins in the bloodstream, releasing GOC in a sustained manner to increase $T_{1/2}$ (Figure 3).

The pharmacokinetic analysis results motivated an exploration into whether OSC-GCDI(P)/HPMCP could extend the dosing interval of the antiviral agent (Figure 5). We orally administrated an identical dose, 10 mg/kg, of OSC-GCDI(P)/HPMCP, as well as OS-P or GOC·TFA, to mice before maPR8 infection at an MLD_{50} of 10. Subsequently, they were treated every day (group 1) or every other day (group 2) for 5 days. To impose more stringent conditions, additional treatments were confined on day 3 (group 3) or day 4 (group 4) or discontinued (group 5) after the first administration (Figure 5a).

When administered with OS-P, there were minimal body weight changes in group 1 with complete survival (Figures 5b and S10a). In groups 2, 3, and 4, body weights initially decreased but have recovered by day 9, resulting in survival rates of 100, 60, and 40%, respectively. However, all mice in group 5 were dead by day 10 after infection. The results well aligned with the short half-life of OS-P less than 1 day.⁵³ GOC·TFA showed antiviral responses comparable or slightly better than OS-P, resulting in complete survival in groups 1, 2, and 3, but 60 and 0% in groups 4 and 5, respectively (Figures 5c and S10b). Notably, when OSC-GCDI(P)/HPMCP was administered, mice survived in all groups (Figures 5d and S10c). Although body weight decreased transiently at the initial stage in group 5, it returned to nearly normal by day 14 postinfection. The data strongly suggested that consistent plasma concentration and extended retention time of GOC metabolized from OSC-GCDI(P) are further enhanced when formulated with HPMCP, making it orally available even after a single shot to treat influenza viral infection.

When OSC-GCDI(P)/HPMCP was administered twice-a-day (b.i.d.) for over 5 days, similar to the standard dosing of OS-P in the clinic, OSC-GCDI(P)/HPMCP did not show the antiviral potency over those of GOC·TFA and OS-P (Figures S8 and S11). The results also supported that much stronger therapeutic effect of OSC-GCDI(P)/HPMCP than GOC·TFA and OS-P with a single oral administration was originated from the maintenance of the GOC concentration above the EC_{50} value for a prolonged period.

In Vivo Antiviral Activity of OSC-GCDI(P)/HPMCP against OS-Resistant Virus. We further explored whether OSC-GCDI(P)/HPMCP is effective against an OS-resistant influenza virus carrying an H275Y mutation, rgKR09, in a mouse model. Following the regimen illustrated in Figure 5a, mice infected with an MLD_{50} of 5 of the virus were orally

administered with OS-P, GOC·TFA, and OSC-GCDI(P)/HPMCP.

When administered with OS-P, both daily treatment (group 1) and every-other-day treatment (group 2) for 5 days exhibited drastic body weight loss and a survival rate of 20%, which was not statistically significant compared to the virus-only group (A/H1N1(H275Y)) (Figures 5e and S12a). In other groups (groups 3–5), it failed to improve body weight changes and survival rates. In another control set with GOC·TFA, all mice were eventually dead, though statistical significance was found in group 1 with a delayed mean survival date (Figures 5f and S12b).

In contrast, when the mice were treated with OSC-GCDI(P)/HPMCP, the alleviation of body weight decrease was particularly evident in group 1 (Figures 5g and S12c). As a result, survival rates improved to 80% in group 1, 40% in group 2, and 20% in groups 3 to 5, demonstrating statistical significance. Although the complete therapeutic efficacy of OSC-GCDI(P)/HPMCP with reduced treatment intervals was not achieved, presumably due to the use of a highly lethal virus dose, therapeutic effectiveness of OSC-GCDI(P)/HPMCP against the OS-resistant virus surpassed that of OS-P and GOC·TFA. Taken together, the results suggest that oral administration of OSC-GCDI(P)/HPMCP not only offers protection against a wild-type influenza virus but also against an OS-resistant variant with NA(H275Y).

CONCLUSION

In the present study, we have developed novel GCDI-based prodrugs of GOC with high oral bioavailability and extended pharmacokinetics, targeting both wild-type and OS-resistant influenza virus strains. The charge of the guanidine group is effectively and temporarily concealed by the GCDI group, increasing the lipophilicity and facilitating the drug absorption through the intestinal barrier. This is the first study of GCDI-based prodrugs with moderate reactivity with nucleophiles, allowing them to be covalently conjugated to serum proteins via a biodegradable linker. GOC, the active compound, can be regenerated in serum in a sustained manner through the hydrolysis of the linker.

Leveraging synergetic effects on intestinal adsorption and slow metabolism, OSC-GCDI(P) has significantly enhanced bioavailability and pharmacokinetic half-life when administered orally. Furthermore, OSC-GCDI(P) showed markedly stronger therapeutic effects compared to OS-P and GOC against both wild-type and H275Y mutant influenza virus infections with just a single oral dose. OSC-GCDIs hold potential as a promising new option against emerging or re-emerging drug-resistant influenza viruses. Additionally, the GCDI-based prodrug design strategy represents a versatile platform technology to enhance the oral bioavailability and serum longevity of various drug candidates containing guanidine groups.

ASSOCIATED CONTENT

Supporting Information

The Supporting Information is available free of charge on the ACS Publications Web site. The Supporting Information is available free of charge at <https://pubs.acs.org/doi/10.1021/acscentsci.4c00548>.

Stability and activation profiles of OSC-GCDIs, proposed activation mechanism of OSC-GCDIs, analysis

data of the OSC-GCDI-HSA conjugates, pharmacokinetic properties, therapeutic efficacy, HPMCP particle characterization, detailed experimental procedures, synthetic procedures, NMR spectra of compounds, and supplemental references (PDF)

Transparent Peer Review report available (PDF)

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Notes

The authors declare no competing financial interest.

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